

John 16188  
#19  
R. Chapman  
3/24/99

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

CONTINUED PROSECUTION APPLICATION (CPA)  
REQUEST TRANSMITTAL

*Priority for Continuation or Divisional applications under 37 CFR 1.53(d)*



Assistant Commissioner for Patents  
Washington, D.C. 20231  
BOX: CPA

Docket No.: 19603/468 (CRF D-1595C)  
Express Mail No.: EL139239885US  
Examiner: J.W. Ricigliano  
Art Unit: 1648

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GROUP 180

This is a Request for filing a:

- continuation or  
 divisional application

under 37 CFR 1.53(d), continued prosecution application, (CPA) of prior application number 08/794,851, filed on February 4, 1997, of Barany et al. entitled **DETECTION OF NUCLEIC ACID SEQUENCE DIFFERENCES USING THE LIGASE DETECTION REACTION WITH ADDRESSABLE ARRAYS**.

**The above-identified prior pending application is hereby expressly abandoned as of the filing date of this request for a continued prosecution application (CPA).**

1.  Enter the unentered amendment previously filed on [\*enter date] under 37 CFR 1.116 in the prior nonprovisional application.
2.  A preliminary amendment is enclosed.
3.  This application is filed by fewer than all the inventors named in the prior application, 37 CFR 1.53 (d)(4).
  - a.  **Delete** the following inventor(s) named in the prior nonprovisional application:

03/04/1999 DURATION 00000096 08794851

01 FC:131  
02 FC:103

760.00 OP  
1242.00 OP

- b.  The inventor(s) to be deleted are set forth on a separate sheet enclosed herewith.
4.  A new power of attorney is enclosed herewith.
5.  Information Disclosure Statement (IDS) with PTO-1449 form(s) and copies of references are enclosed herewith.

03/24/1999 R.HARMON 00000005 161138 08794851  
01 FC:103 6. The filing fee is calculated on the basis of the claims existing in the prior application as  
180.00 intended above:

(Col. 1)		(Col. 2)		SMALL ENTITY		LARGE ENTITY		
FOR:	NO. FILED	NO. EXTRA	RATE	FEE	OR	RATE	FEE	
BASIC FEE	XXXXXXXXXX	XXXXXXXXXX	XXXXXX	\$380	OR	XXXXX	\$760	
TOTAL CLAIMS	89 - 20 =	69	x 9 =	\$	OR	x 18 =	\$1242	
INDEP CLAIMS	1 - 3 =	0	x 39 =	\$	OR	x 78 =	\$	
[ ] MULTIPLE DEPENDENT CLAIM PRESENTED				x130 =	\$	OR	x260 =	\$
				TOTAL	\$	OR	TOTAL	\$2002

\*If the Total Claims are less than 20 and Indep. Claims are less than 3, enter "0" in Col. 2

7.  A check in the amount of \$2,002.00 is enclosed to cover the above filing fee.
8.  The Commissioner is hereby authorized to charge fees which may be required, or credit overpayment to Deposit Account No. 14-1138. A duplicate copy of this sheet is enclosed.
9. Small entity status:
  - A small entity statement is enclosed.
  - A small entity statement was filed in the prior nonprovisional application and such status is still proper and desired.
  - is no longer claimed.
10.  A copy of the Request for One-Month Extension of Time in prior application.
11.  A self-addressed, prepaid postcard for acknowledging receipt.
12. Other:

Address all future communications to (may only be completed by Applicant, or attorney or agent of record):

Michael L. Goldman  
Nixon, Hargrave, Devans & Doyle LLP  
Clinton Square  
P.O. Box 1051  
Rochester, New York 14603

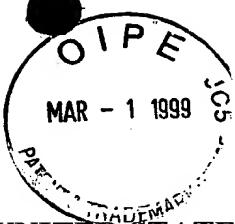
Respectfully submitted,



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**Docket No.: 19603/468 (CRF D-1595C)**

*J. P. D.*  
J. P. D.  
PATENT  
F D-1595C)  
E 3/24/59

**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

Applicants	:	Barany et al.	)	Examiner:
Serial No.	:	CPA of 08/794,851	)	J.W. Ricigliano
Filed	:	February 4, 1997	)	Art Unit: 1648
For	:	DETECTION OF NUCLEIC ACID SEQUENCE DIFFERENCES USING THE LIGASE DETECTION REACTION WITH ADDRESSABLE ARRAYS	)	

## **PRELIMINARY AMENDMENT**

Assistant Commissioner for Patents  
Washington, D.C. 20231 -

Dear Sir:

Please cancel claims 89-137 without prejudice.

Please amend the above-identified patent application as follows:

**In the Claims:**

Please add new claim 148 as follows:

*84* 48. A method according to claim 1, wherein the capture oligonucleotides are in the form of DNA or PNA.

Please amend claims 1, 44, and 81 as follows:

1. (Amended) A method for identifying one or more of a plurality of sequences differing by one or more single-base changes, insertions, deletions, or translocations in a plurality of target nucleotide sequences comprising: